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GenCore version 5.1.6
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OM protein - protein search, using sw model

May 31, 2003, 17:57:32 ; Search time 15.6444 Seconds Run on:

(without alignments) 828.194 Million cell updates/sec

US-09-772-103-8

1 MDFQVQIFSFLLISASVILS......CQQWSSYPLTFGGGTKVEIK 128 Perfect score: Sequence:

BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

383519 seqs, 101223694 residues ched:

'number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Published Applications AA:*

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2: /cgn2_6/ptodata/1/pubpaa/PCT NEW PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/US06_NEW PUB.pep:*
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6: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
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8: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description		Sequence 92, Appl	Sequence 52, Appl	Sequence 99, Appl	Sequence 94, Appl	Seguence 97 April	• "	הקקה יפי המוניה אלים		sednence II3, App		Sequence 93, Appl			Sequence 50, Appl	Sequence 61, Appl	Sequence 71, Appl	Seguence 87. April		0	4,	Sequence 6. Appli	ហ	
	ID	TIS-10-140-222-02	TIS-00-010-05	75-05-016-60-52	US-US-ATU-USA-AA	US-IU-I6U-Z3Z-94	US-09-910-059-97	US-10-160-232-96	US-10-195-752-111	175-10-195-752-112	TIO 010 010 11	/T-650-0T6-60-60	US-10-160-232-93	US-09-753-436-96	113-09-910-010-01	TIS -010-010-00	T9-650-016-60-65	T/-690-0T6-60-90	US-10-160-232-87	US-10-160-232-89	TIS 00 - 0 - 0 - 0 - 0 - 0 - 0 - 0 - 0 - 0	110 00 000 010 0	02-03-36/-/3C-6	US-09-795-515-5	
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9 US-09-771-415-1 10 US-09-796-848A 9 US-09-996-288-2 9 US-09-996-265-2			9 US-09-905-928 9 US-10-096-964 10 US-10-95-09 12 US-10-051-85, 9 US-09-996-288 9 US-09-996-288	9 US-09-996-288-24 9 US-09-996-265-23 9 US-09-996-265-23 9 US-09-996-265-23
4.47 4.47 4.47 4.47 4.47 4.47 4.47		482 73.6 482 73.6 481 73.4 481 73.4 481 73.4	481 73.4 481 73.4 481 73.4 481 73.4 481 73.4	
	4 7 0 4 8 6	6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	3 3 3 3 3 4 4 4 4 4 0 0 1 4 4 1 1 4 4 1 1 1 1 1 1	54 4 4 4 5 4 5 5

ALIGNMENTS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: Synthetic protein
                                                                                                                                                  APPLICANT: ITO, MIKITO
APPLICANT: HANAI, NOBUO
APPLICANT: KAWADA, VOKO
APPLICANT: NAKAMURA, KAZUYASU
APPLICANT: NAKAMURA, KAZUYASU
APPLICANT: STBUTA, MASABUMI
TITLE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 91.4%; Pred. No. 2.7e-36;
Matches 117; Conservative 5; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2002-06-04
PRIOR APPLICATION UNDER: US/09/453,718
PRIOR PILING DATE: 1999-12-03
PRIOR PLILNG DATE: 1999-05-20
PRIOR FILING DATE: 1999-05-20
PRIOR PLICATION NUMBER: 09/119,014
PRIOR APPLICATION NUMBER: 09/119,014
PRIOR PLILNG DATE: 1998-07-20
PRIOR PLING DATE: 1998-07-20
PRIOR FILING DATE: 1997-11-21
NUMBER OF SEQ ID NOS: 96
SOFTWARE: PATCHILL VET: 2.1
                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/160,232 CURRENT FILING DATE: 2002-06-04
                                                 ; Sequence 92, Application US/10160232; Publication No. US20030088075A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                           APPLICANT: SHITARA, KENYA
RESULT 1
US-10-160-232-92
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61 PGKAPKLLIYDTSNLASGVPSRFSGSGSGTDYTLTISSLQPEDFATYYCQQMSSYPLTFG 120

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61 PGKAPKLLIYDTSNLASGVPSRFSGSGSGTDYTLTISSLQPEDFATYXCQQWSSYPLTFG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 PGKAPKLMIYSTSNLASGVPSRFSGSGSGTDYTFTISSLOPEDIATYYCQQRSTYPLTFG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: INITATIO, MIKITO
APPLICANT: ITO, MIKITO
APPLICANT: HANAI, NOBUO
APPLICANT: KAMADA, YOKO
APPLICANT: KAMADA, YOKO
APPLICANT: SHIBUYA, MASABUMI
APPLICANT: SHIBUYA, MASABUMI
APPLICANT: SHIBUYA, MASABUMI
TITUE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY
FILE REFERENCE: 249-107
CURRENT APPLICATION NUMBER: US/10/160,232
CURRENT FILING DATE: 1999-12-03
PRIOR FILING DATE: 1999-12-03
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1999-07-20
PRIOR FILING DATE: 1997-11-21
PRIOR SEQIID NOWERS: PCT/JP97/04259
PRIOR FILING DATE: 1997-11-21
NUMBER OF SEQIID NOS: 96
                                                                                                                                                                                                                                                                                                                                                                                                                                         1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Synthetic protein
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                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
                                                                                                                                                                                                                                                                                                                                            90.5%; Score 593; DB 10; Length 235; 90.6%; Pred. No. 1.5e-35; tive 5; Mismatches 7; Indels C
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                                                                                                                                                                                                                                                                      , OTHER INFORMATION: humanised light chain Fd sequence US-08-910-059-99
PRIOR APPLICATION NUMBER: GB 9703103.3 PRIOR FILING DATE: 1997-02-14 PRIOR APPLICATION NUMBER: GB9609405.7 PRIOR FILING DATE: 1996-05-04 NUMBER OF SEQ ID NOS: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 94, Application US/10160232; Publication No. US20030088075A1; GENERAL INFORMATION:
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Matches 116; Conservative
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Best Local Similarity
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SEQ ID NO 94
LENGTH: 126
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                                                                                                                                                              SEQ ID NO 99
LENGTH: 235
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APPLICANT: Copley, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Edge, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
TITLE OF INVENTION: Their Therapeutic use in an Adept System
                                                                                                                                                                                                                                                                                                                               APPLICANT: COLLEY, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
TITLE OF INVENTION: Their Therapeutic use in an Adept System
FILE REFERENCE: 1991-209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 PGKAPKLLIYDISNLASGVPSRFSGSGGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
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91.5%; Score 599; DB 10; Length 235;
Best Local Similarity 91.4%; Pred. No. 5.5e-36;
Matches 117; Conservative 5; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: complete humanised light chain sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TRIBLY APPLICATION NUMBER: US/09/910,059

(TRENT FILING DATE: 2001-07-23

PRIOR APPLICATION NUMBER: US 09/171,945

PRIOR APPLICATION NUMBER: US 09/171,945

PRIOR PILING DATE: 1998-10-29

PRIOR PILING DATE: 1997-04-29

PRIOR PILING DATE: 1997-04-29

PRIOR PILING DATE: 1997-02-14

PRIOR PILING DATE: 1997-02-14

PRIOR PILING DATE: 1996-05-04

NUMBER OF SEQ ID NOS: 131

SOFTWARE: PATENTIN VERSION 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 1991-209
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT FILING DATE: 2001-07-23
PRIOR YAPLICATION NUMBER: US 09/171,945
PRIOR FILING DATE: 1998-10-29
PRIOR PLING DATE: 1998-10-29
PRIOR PLING DATE: 1997-04-29
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Patent No. US20020142359A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                      ; Sequence 52, Application US/09910059; Patent No. US20020142359A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                               RESULT 2
US-09-910-059-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 52
LENGTH: 235
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61 PGKAPKLLIYDTSNLASGVPSRFSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
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                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic protein US-10-160-232-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 87.5%; Score 573; DB 9; Length 126; Best Local Similarity 89.1%; Pred. No. 2.3e-34; Matches 114; Conservative 5; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: AKLINGLON
COUNTRY: U.S.A.

ZIP: 22201-4714
COMPUTER READBELE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/195,752
FILING DATE: 16-Jul-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION AURAER: US/09/393,385B
FILING DATE: 27-JUN-96
TELECOMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KUWANA, YOSHIHISA
HASEGAWA, MAMORU
TITLE OF INVENTION: HUMANIZED ANTIBODIES
NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: NIXON & VANDERHYE P.C. STREET: 1100 NORTH GLEBE ROAD
             PRIOR APPLICATION NUMBER: 09/315,051
PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 09/119,014
PRIOR PLING DATE: 1998-07-20
PRIOR APPLICATION NUMBER: PCT/JP97/04259
PRIOR FILING DATE: 1997-11-21
NUMBER OF SEQ ID NOS: 96
SEQ ID NO 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 111, Application US/10195752
Publication No. US20030077276A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (703)816-4000
TELEFAX: (703)816-4100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 130 amino acids
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SHITARA, KENYA
HANAI, NOBUO
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SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                               LENGTH: 126
                                                                                                                                                                                                                                                                       TYPE: PRT
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                                                                                                                                                                                                                                               General No. 91, Application US/09910059

Fatent No. US20020142359A1

GENERAL INFORMATION:

APPLICANT: COPIEY, Clive G

APPLICANT: Edge, Michael Derek

APPLICANT: Edge, Michael Derek

APPLICANT: Edge, Michael Antibody to CEA, Conjugates Comprising Said Antibody,

TITLE OF INVENTION: Their Therapeutic use in an Adept System

TITLE OF INVENTION: Their Therapeutic use in an Adept System

TITLE OF INVENTION: Their Therapeutic use in an Adept System

FILE REFERENCE: 1991-209

REBUT APPLICATION NUMBER: US/09/910,059

PRIOR PILING DATE: 1998-10-29

PRIOR PILING DATE: 1998-10-29

PRIOR PILING DATE: 1997-04-29

PRIOR APPLICATION NUMBER: GB 9703103.3

PRIOR FILING DATE: 1997-04-29

PRIOR FILING DATE: 1997-05-04

NUMBER OF SEQ ID NOS: 131

SEQ ID NO 97

TAPP: DEPT.

TAPP: DEPT.

TAPP: DEPT.

TAPP: DEPT.

TAPP: DEPT.
61 PGKAPKLLIYRTSNLASGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCHQWSMY--TFG 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 PGKAPKLLIYDTSNLASGVPSRFSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
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Publication No. US20030088075A1
GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBHO
APPLICANT: SHIBADA, WORK
APPLICANT: SHIBADA, WORK
APPLICANT: SHIBADA, WORK
APPLICANT: SHIBUYA, MASABUMI
TITLE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY
FILE REFERENCE: 249-107
CURRENT APPLICATION NUMBER: US/10/160,232
CURRENT FILING DATE: 2002-06-04
PRIOR APPLICATION NUMBER: US/09/453,718
PRIOR FILING DATE: 1999-12-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK
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89.1%; Pred. No. 7.6e-35;
tive 6; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: humanised light chain Fd sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 89.18
Matches 114; Conservative
                                                                               121 GGTKVEIK 128
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121 QGTKLEIK 128
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PGKAPKLLIYDTSNLASGVPSRFSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
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85.9%; Pred. No. 1e-33;
iive 8; Mismatches 10; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: MISCANAL OF AN ACCOUNTRY: U.S.A.

ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/195,752
FILING DATE: 16-Jul-2002
CLASSIFICATION: <UNKNOWN>
                                                                                                        Length 130;
                                                                                                                                                 10; Indels
                                                                                                   Score 567; DB 9;
Pred. No. 6.2e-34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KUWANA, YOSHIHISA
HASEGANA, WAMORU
TITLE OF INVENTION: HUMANIZED ANTIBODIES
NUMBER OF SEQUENCES: 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/393,385B
FILING DATE: 27-JUN-96
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)816-4000
TELEPHONE: (703)816-4100
                                                                                                                                                 7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSER: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 113:
US-10-195-752-113
      MOLECULE TYPE: process

) SEQUENCE DESCRIPTION: SEQ ID NO: 111:

US-10-195-752-111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 113, Application US/10195752
Publication No. US20030077276A1
GENERAL INFORMATION:
APPLICANT: NAKAMURA, KAZUYASU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 130 amino acids TYPE: amino acid
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SHITARA, KENYA
HANAI, NOBUO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 113:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single
                                                                                                   Query Match
Best Local Similarity 86.7%;
Aatches 111; Conservative
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Best Local Similarity 85.9°
Matches' 110; Conservative
TOPOLOGY: linear
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                                                                                                          Query Match
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1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK 60

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GREEAL INCRMATION:

GREEAL INCRMATION:

GREEAL INCRMATION:

GREEAL SERVER CONTINUED CHARLES

APPLICANT: Edge, Michael Derek

APPLICANT: Edge, Michael Derek

TTPLE OF INVENTION: Their Therapeutic use in an Adept System

TITLE OF INVENTION: Their Therapeutic use in an Adept System

TITLE OF INVENTION: Their Therapeutic use in an Adept System

TITLE OF INVENTION: Their Therapeutic use in an Adept System

TITLE OF INVENTION: Their Therapeutic use in an Adept System

TITLE OF INVENTION: Their Therapeutic use in an Adept System

TITLE OF INVENTION: Their Therapeutic use in an Adept System

TITLE OF INVENTION: 1991-0.23

CURRENT APPLICATION NUMBER: US 09/171,945

PRIOR FILING DATE: 1997-04-29

PRIOR APPLICATION NUMBER: GB 9703103.3

PRIOR PRILING DATE: 1997-02-14

PRIOR FILING DATE: 1996-05-04

NUMBER OF SEQ ID NOS: 131

SEQ ID NO 17

LENGTH: 235
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                                                           61 PGKAPKILIYDTSNLASGVPSRFSGSGGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
                                                                                         61 PGKAPKIMIYSTSNLASGVPARFSGSGSGTSYSLTISRLQPEDIATYYCQQRSSYPYTFG 120
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APPLICANT: NAKAMURA, KAZIYASU
APPLICANT: NHINYA, MASABUMI
ITILE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY
FILE REFERENCE: 249-107
CURRENT APPLICATION NUMBER: US/10/160,232
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Pest Local Similarity 77.3%; Pred. No. 1.5e-30;
Matches 99; Conservative 17; Mismatches 12;
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                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 17, Application US/09910059; Patent No. US20020142359A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            121 GGTKVEIK 128
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61 PGKAPKLLIYDTSNLASGVPSRFSGSGGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK 60
                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: Synthetic protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     itch     79.4%; Score 520; DB 9; Length 126;
cal Similarity 79.7%; Pred. No. 1.4e-30;
102; Conservative 9; Mismatches 15; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gallatin, W. Michael
APPLICANT: Vazeux, Rosemay
TTLE OF INVENTION: ICAM-Related Materials and Methods
NUMBER OF SEQUENCES: 120
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun
STREET: 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 6300 Saars Tower, 233 South Wacker Drive CITY: Chicago Saars Tower, 233 South Wacker Drive STATE: 111inois
COUNTRY: United States of America ZIP: 60606-6402
COUNTRY: United States of America ZIP: 60606-6402
COMPUTER READABLE FORM:
MEDIUM TYPE: FORM:
MEDIUM TYPE: FISH PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/753,436
CURRENT FILING DATE: 2002-06-04

PRIOR APPLICATION NUMBER: US/09/453,718

PRIOR FILING DATE: 1999-12-03

PRIOR APPLICATION NUMBER: 09/315,051

PRIOR APPLICATION NUMBER: 09/119,014

PRIOR PILING DATE: 1999-05-20

PRIOR PILING DATE: 1998-07-20

PRIOR PLING DATE: 1998-07-20

PRIOR FILING DATE: 1997-11-21

PRIOR FILING DATE: 1997-11-21

SEQ ID NOS: 96

SOFTWARE: PATENTIN Ver: 2.1

SEQ ID NO 93

LING APPLICATION VARIABLES

SEQ ID NO 93
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/286,754
FILING DATE: 05-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/102,852
FILING DATE: 05-AUG-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/487,113
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/382,289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 96, Application US/09753436; Patent No. US20010029293A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       121 GGTKVEIK 128
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US-09-753-436-96
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Sequence 50, Application US/09910059
; Sequence 50, Application US/09910059
; Patent No. US2002014235941
; GENERAL INFORMATION:
    APPLICANT: Copiley, Clive G
    APPLICANT: Edge, Michael Derek
    TILE OF INVENITON: Their Therapeutic use in an Adept System
    TILING DATE: 1991-20
    CURRENT FILING DATE: 1998-10-29
    PRIOR PILING DATE: 1999-10-29
    PRIOR PILING DATE: 1997-04-29
    PRIOR PILING DATE: 1997-04-29
    PRIOR PILING DATE: 1997-02-14
    PRIOR PILING DATE: 1996-05-04
    PRIOR PILING DATE: 1996-05-04
    NUMBER OF SEQ ID NOS: 131
    SOFTWARE: PatentIn version 3.1
    SOFTWARE: PatentIn version 3.1
    LENGTH: 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 SRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQKPGKAPKLLIYDTSNLASGV 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 ARCDIQMTQSPSSLSASVGDRVTITCSASSSVSYIYWYQQKPGKAPKLLIYLTSNLASGV 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          62 PSRFSGSGSGTDFTLTISSLQPEDFATYYCQQWKSIPLTFGQGTKLEIK 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    76.3%; Score 500; DB 10; Length 110;
88.1%; Pred. No. 3.3e-29;
tive 7; Mismatches 6; Indels (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               76.2%; Score 499; DB 10; 90.6%; Pred. No. 3.8e-29;
               FILING DATE: 20-JAN-1993
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 05-UN1-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/889,724
FILING DATE: 26-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/827,689
FILING DATE: 21-DAN-1992
ATTORNEY/AGENT INFORMATION:
NAME: Williams, JOSEPA A., Jr.
REGISTRATION NUMBER: 33,282
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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TELEFAX: (312) 474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 96:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            110 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 76.3%
Best Local Similarity 88.1%
Matches 96; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear; MOLECULE TYPE: protein US-09-753-436-96
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
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Best Local Similarity
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US-09-910-059-50
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TYPE: PRT
ORGANISM: Artificial Sequence
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11 Similarity 89.6%;
95; Conservative
                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: ITO, MIKITO
APPLICANT: HANAI, NOBUO
APPLICANT: KAWADA, YOKO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     121 GGTKVEIK 128
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                                                                                                                                     LENGTH: 107
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Best Local S
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GENERAL INFORMATION:

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Edge, Michael Derek

APPLICANT: Edge, Michael Derek

APPLICANT: Emery. Stephen Charles

TILE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,

TEATLE OF INVENTION: Their Therapeutic use in an Adept System

LE REFERENCE: 1991-209

CURRENT APPLICATION NUMBER: US/09/910,059

CURRENT APPLICATION NUMBER: US 09/171,945

PRIOR APPLICATION NUMBER: PCT/GB97/01165

PRIOR FILING DATE: 1997-02-14

PRIOR FILING DATE: 1997-02-14

PRIOR FILING DATE: 1997-02-14

PRIOR FILING DATE: 1996-05-04

NUMBER OF SEQ ID NOS: 131

SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 11, Application US/09910059
; Patent No. US20020142359A1
; General INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Copley, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
TITLE OF INVENTION: Their Therapeutic use in an Adept System
FILE REFERENCE: 1991-209
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT APPLICATION NUMBER: US 09/171,945
PRIOR *TILING DATE: 1998-10-29
; PRIOR *TILING DATE: 1998-10-29
; PRIOR *FILING DATE: 1997-04-29
; PRIOR FILING DATE: 1997-04-29
; PRIOR FILING DATE: 1997-04-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 DIQLIQSPSSISASVGDRVIITCSASSSVIYMHWYQQKPGKAPKLLIYSTSNLASGVPSR 60
                                                                                        1 DIOMIQSPSSLSASVGDRVIITCSASSSVTYMHWYQQKPGKAPKLLIYSTSNLASGVPSR 60
                                             DIOMIQSPSSLSASVGDRVTITCSATSSITYMSWYQQKPGKAPKLLIYDTSNLASGVPSR 82
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0; Gaps
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; OTHER INFORMATION: humanised light chain variable region variant
V3-09-910-059-61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          83 FSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFGGGTKVEIK 128
                                                                                                                                             83 FSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFGGGTKVEIK 128
                                                                                                                                                                   6; Indels
  4; Mismatches
                                                                                                                                                                                                                                                                                                                 Sequence 61, Application US/09910059 Patent No. US20020142359A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
  96; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                          US-09-910-059-61
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LENGTH: 107
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     Matches
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61 PGKAPKILIYDTSNIASGVPSRFSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23 DIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQKPGKAPKLLIYDTSNLASGVPSR
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APPLICANT: SHIBUYA, MASABUMI
TITLE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY
FILE REFERENCE: 249-107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , OTHER INFORMATION: humanised light chain variable region variant US-09-910-059-71
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 493; DB 10; Length 107; Pred. No. 1e-28; 7; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 75.3%; Score 493; DB 9; Length 128; Best Local Similarity 72.7%; Pred. No. 1.2e-28; Matches 93; Conservative 18; Mismatches 17; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              83 FSGSGSGTDYTLTISSLOPEDFATYYCQQWSSYPLTFGGGTKVEIK 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Mismatches
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PRIOR FILING DATE: 1997-02-14
PRIOR APPLICATION NUMBER: GB9603405.7
PRIOR FILING DATE: 1996-05-04
NUMBER OF SEQ ID NOS: 131
SOFTWARE: PatentIn version 3.1
SEQ ID NO 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 87, Application US/10160232 Publication No. US20030088075A1 GENERAL INFORMATION: APPLICANT: SHITARA, KENYA
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Page 7

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Search completed: May 31, 2003, 18:05:42 Job time : 17.6444 secs

